WO 00/23601

PCT/AU99/00896

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Sequence listing:

Applicants: Commonwealth Scientific and Industrial Research Organisation

University of Western Sydney (Nepean)
Pig Research and Development Corporation

Title of the Invention: Delivery system for porcine somatotropin

10

Prior Application Number: PP 6556

Prior Application Filing Date: 1998-10-16

Number of SEQ ID NOs: 3

15

25

Software: PatentIn Ver. 2.1

SEQ ID NO: 1

Length: 72

20 Type: DNA

Organism: Homo sapien

Sequence: 1

atggccctgt ggatgcgcct cctgcccctg ctggcgctgc tggccctctg gggacctgac 60

ccagccgcag cc

SEQ ID NO: 2

Length: 666

30 Type: DNA

Organism: Artificial Sequence

Feature:

Other Information: Description of Artificial Sequence: ISS-pST gene

35 construct



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Sequence: 2 gctagcatgg ccctgtggat gcgcctcctg cccctgctgg cgctgctggc cctctgggga 60 cctgacccag ccgcagccct cgagatgttt ccagctatgc cactttcttc tctgttcgct 120 aacgctgttc ttcgggccca gcacctgcac caactggctg ccgacaccta caaggagttt 180 5 gagegegeet acatecegga gggacagagg tactecatee agaaegeeea ggetgeette 240 tgcttctcgg agaccatccc ggccccacg ggcaaggacg aggcccagca gagatcggac 300 gtggagetge tgegettete getgetgete atceagtegt ggetegggee egtgeagtte 360 ctcagcaggg tcttcaccaa cagcctggtg tttggcacct cagaccgcgt ctacgagaag 420 ctgaaggacc tggaggaggg catccaggcc ctgatgcggg agctggagga tggcagcccc 480 10 cgggcaggac agatcctcaa gcaaacctac gacaaatttg acacaaactt gcgcagtgat 540 gacgcgctgc ttaagaacta cgggctgctc tcctgcttca agaaggacct gcacaaggct 600 gagacatace tgegggteat gaagtgtege egettegtgg agageagetg tgeettetag 660 666 tctaga 15 SEQ ID NO: Length: 217 Type: PRT 20 Organism: Artificial Sequence Feature: Other Information: Description of Artificial Sequence: ISS-pST peptide sequence 25 Zp. . Sequence: Met Ala Leu Trp Met Arg Leu Leu Pro Leu Leu Ala Leu Leu Ala Leu 15 10 1 Trp Gly Pro Asp Pro Ala Ala Ala Leu Glu Met Pne Pro Ala Met Pro 30 30 20 25

Leu Ser Ser Leu Phe Ala Asn Ala Val Leu Arg Ala Gln His Leu His 40

35

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Gln Leu Ala Ala Asp Thr Tyr Lys Glu Phe Glu Arg Ala Tyr Ile Pro

Glu Gly Gln Arg Tyr Ser Ile Gln Asn Ala Gln Ala Ala Phe Cys Phe

Ser Glu Thr Ile Pro Ala Pro Thr Gly Lys Asp Glu Ala Gln Gln Arg

Ser Asp Val Glu Leu Leu Arg Phe Ser Leu Leu Ile Gln Ser Trp

Leu Gly Pro Val Gln Phe Leu Ser Arg Val Phe Thr Asn Ser Leu Val

Phe Gly Thr Ser Asp Arg Val Tyr Glu Lys Leu Lys Asp Leu Glu Glu

Gly Ile Gln Ala Leu Met Arg Glu Leu Glu Asp Gly Ser Pro Arg Ala

Gly Gln Ile Leu Lys Gln Thr Tyr Asp Lys Phe Asp Thr Asn Leu Arg

Ser Asp Asp Ala Leu Leu Lys Asn Tyr Gly Leu Leu Ser Cys Phe Lys

Lys Asp Leu His Lys Ala Glu Thr Tyr Leu Arg Val Met Lys Cys Arg

Arg Phe Val Glu Ser Ser Cys Ala Phe